IN THE SPECIFICATION

Please amend the specification as shown:

Please delete Table 2 on pages 14-20 and replace it with the following table:

Table 2 (FRH1-4 disclosed as SEQ ID NO: 151 and FRL1-4 disclosed as SEQ ID NO: 152):

| Name | Position range | Pos. | Amino acid or |
|------|--------------------------|------|---------------------|
| | | | amino acid sequence |
| FRH1 | 1 to 30 (SEQ ID NO: 143) | 1 | Q or E |
| | | 2 | V |
| | | 3 | Q, K or T |
| | | 4 | L |
| | | 5 | KorV |
| | | 6 | E or Q |
| | | 7 | S |
| | | 8 | G |
| | | 9 | A |
| | | 10 | E |
| | | 11 | L or V |
| | | 12 | V or K |
| | | 13 | R or K |
| | | 14 | Р |
| | | 15 | G |
| | | 16 | T or A |
| | | 17 | S |
| | | 18 | V |
| | | 19 | K |
| | | 20 | I or V |
| | | 21 | S or P |
| | | 22 | С |
| | | 23 | K |
| | | 24 | A, V, S or T |
| | | 25 | S |
| | | 26 | G |

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| | | 27 | Y, F, S or D |
|-------|---------------------------|----|---------------------------|
| | | 28 | Т |
| | | 29 | F, L or I |
| | | 30 | Т |
| CDRH1 | 31 to 35 | | SEQ ID NO. 1 and variants |
| FRH2 | 36 to 49 (SEQ ID NO: 144) | 36 | W |
| · · · | | 37 | V |
| | | 38 | K or R |
| | | 39 | Q |
| | | 40 | R or A |
| | | 41 | Р |
| | | 42 | G |
| | | 43 | H or Q |
| | | 44 | G |
| | | 45 | L |
| | | 46 | E |
| | | 47 | W or R |
| | | 48 | I or M |
| | | 49 | G |

| 50 to 65, with position 52a | | SEQ ID NO. 2 or 3 and |
|---------------------------------------|------------------------|---|
| introduced in addition | | variants |
| 66 to 94 (SEQ ID NO: 145) | 66 | K or R |
| | 67 | A or V |
| | 68 | Т |
| | 69 | L or M |
| | 70 | Т |
| | 71 | A, L or T |
| | 72 | D |
| | 73 | Т |
| | 74 | S |
| | 75 | S or T |
| | 76 | S |
| | 77 | Т |
| | 78 | Α |
| · · · · · · · · · · · · · · · · · · · | introduced in addition | introduced in addition 66 to 94 (SEQ ID NO: 145) 68 69 70 71 72 73 74 75 76 77 |

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| | | 79 | Y |
|-------|---------------------------------------|-----|--------------------------|
| | | 80 | M |
| | | 81 | Q or E |
| | | 82 | L |
| | | 82a | S |
| | · · · · · · · · · · · · · · · · · · · | | |
| | | 82b | S or R |
| | | 82c | L |
| | | 83 | T or R |
| | | 84 | S |
| | | 85 | E |
| | | 86 | D |
| | | 87 | S or T |
| | | 88 | Α |
| | | 89 | V |
| | | 90 | Y |
| | | 91 | ForY |
| | | 92 | С |
| | | 93 | A |
| | | 94 | Y, K or R |
| CDRH3 | 95 to 102, with positions 100a | | SEQ ID NO. 4, 5 or 6 and |
| | and 100b introduced in addition | | variants |
| FRH4 | 103 to 113 (SEQ ID NO: 146) | 103 | W |
| | | 104 | G |
| | | 105 | Q |
| | | 106 | G |
| | | 107 | T |
| | | 108 | T, S or L |
| | | 109 | V or L |
| | | 110 | T |
| | | 111 | V |
| | | 112 | S |
| | | 113 | S or A |
| FRL1 | 1 to 23 (SEQ ID NO: 147) | 1 | D |
| | | 2 | I, V or L |
| | | 3 | |
| | <u> </u> | | Q or L |

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| | | 4 | M |
|---------------------------------------|-------------------------------------|-----|--------------------------|
| | | 5 | T |
| | | 6 | Q |
| | | 7 | T or S |
| | | | |
| | | 8 | P |
| | | 9 | L |
| | | 10 | S |
| | | 11 | L |
| | | 12 | Р |
| | | 13 | V |
| | | 14 | S or T |
| | | 15 | L or P |
| | | 16 | G |
| | | 17 | D or E |
| | | 18 | Q or P |
| | | 19 | A |
| | | 20 | S |
| | | 21 | |
| | | 22 | S |
| | | 23 | C |
| CDDI 4 | (100)104 to 04 with a siting 07- | 25 | |
| CDRL1 | [[22]]24 to 34, with positions 27a, | | SEQ ID NO. 7, 8 or 9 and |
| | 27b, 27c, 27d and 27e | | variants |
| | introduced in addition | | |
| FRL2 | 35 to 49 (SEQ ID NO: 148) | 35 | W |
| · · · · · · · · · · · · · · · · · · · | | 36 | Υ |
| | | 37 | L |
| | | 38 | Q |
| , | | 39 | К |
| | | 40 | Р |
| | | 41 | G |
| | | 42 | Q |
| | | 43 | S |
| | | 44 | P |
| | | 45 | K or Q |
| | | 46 | L |
| | | 170 | |

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| | | 47 | L |
|-------|---------------------------|----|----------------------------------|
| | | 48 | I or V |
| ** . | | 49 | Y |
| CDRL2 | 50 to 56 | | SEQ ID NO. 10 or 11 and variants |
| FRL3 | 57 to 88 (SEQ ID NO: 149) | 57 | G |
| | | 58 | V |
| | | 59 | Р |
| | | 60 | D |
| | | 61 | R |
| | | 62 | F |
| • | | 63 | S |
| | | 64 | G |
| | | 65 | S |
| | | 66 | G |
| | | 67 | S |
| | | 68 | G |
| | | 69 | Т |
| | | 70 | D |
| | | 71 | F |
| | | 72 | Т |
| | | 73 | L |
| | | 74 | К |
| | | 75 | I |
| | | 76 | S |
| | | 77 | R |
| | | 78 | V |
| | | 79 | E |
| , | | 80 | A |
| | | 81 | E |
| | | 82 | D |
| | | 83 | L or V |
| , | | 84 | G |
| | | 85 | V |
| | | 86 | Y |

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| | | 87 | Υ |
|-------|----------------------------|------|----------------------------------|
| | | 88 | С |
| CDRL3 | 89 to 97 | | SEQ ID NO. 12 or 13 and variants |
| FRL4 | 98 to 108 (SEQ ID NO: 150) | 98 | F |
| | | 99 | G |
| | | 100 | G or Q |
| | | 101 | G |
| | | 102 | Т |
| | | 103 | К |
| | | 104 | L |
| | | 105 | E |
| | | 106 | l or L |
| | | 106a | К |
| | | 107 | R |
| | | 108 | Α |

Please delete the paragraphs on page 77, lines 13-18 and replace them with the following paragraphs:

Fig. 1a: Sequences of linkers in various multibody single-chain antibody fragments (SEQ ID NOS 114-135, respectively, in order of appearance)

Fig. 1b: Cloning diagram for the preparation of single-chain antibody fragments having different linker length (6 His tag is disclosed as SEQ ID NO: 136).

Please delete the paragraph on page 78, lines 1-3 and replace it with the following paragraph:

Fig. 4: Vector system for cloning and eukaryotic expression of chimeric antibodies in IgG1 or IgM format. <u>Figure discloses SEQ ID NOS 137-142</u>, respectively, in order of appearance.

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